SEQUENCE LISTING

<110> Hamm, Heidi
      Gilchrist, Annette

<120> Method For Identifying Inhibitors of G Protein Coupled Receptor
Signaling

<130> 2661-101

<140> US 09/852,910
<141> 2001-05-11

<150> US 60/275,472
<151> 2001-03-14

<160> 271

<170> PatentIn version 3.0

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1   5   10

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1   5   10

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1   5   10

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1   5   10

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1   5   10

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1   5   10

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1   5   10

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Ile Ile Ser Ala Ser Leu Lys Met Val Gly Val
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Asn Glu Asn Leu Arg Ser Ala Gly Leu His Glu
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1   5   10

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1   5   10

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Ser Gln Val Lys Leu Gln Lys Gly His Leu Val
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Leu Gln Leu Arg Leu Arg Glu His Asn Leu Val
1   5   10

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1   5   10

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1   5   10

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Leu Gln Lys Asn Leu Lys Glu Tyr Asn Met Val
1   5   10

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<400>  107

Leu Gln Tyr Asn Leu Met Glu Asp Tyr Leu Asn
1                5                10


<210>  108
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<400>  108

Leu Gln Met Tyr Leu Arg Gly Tyr Asn Leu Val
1                5                10


<210>  109
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<400>  109

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1                5                10


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<400>  110

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1                5                10


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1   5   10

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Met Gly Ile Lys Asn Asn Leu Lys Asp Cys Gly Leu Phe
1   5   10

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1   5   10

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1   5   10

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1   5   10

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1   5   10

<210> 117
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agtaaagc   68

<210> 119
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                                                                        67
gtaaagc


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gtaaagc   67

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                                                                          67

gtaaagc

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<400> 124

Ile Arg Glu Asn Leu Lys Asp Cys Gly Leu Phe
1   5   10

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<400> 125

Leu Leu Glu Asn Leu Arg Asp Cys Gly Met Phe
1   5   10

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<400> 126

Ile Gln Gly Val Leu Lys Asp Cys Gly Leu Leu
1   5   10

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<400> 127

Ile Cys Glu Asn Leu Lys Glu Cys Gly Leu Phe
1   5   10

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<400> 128

Met Leu Glu Asn Leu Lys Asp Cys Gly Leu Phe
1   5   10

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1   5   10

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1   5   10

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Leu Leu Asp Asn Ile Lys Asp Cys Gly Leu Phe
1   5   10

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1   5   10

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<400> 133

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1   5   10

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<400> 134

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1   5   10

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<400>  135

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1               5                   10


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<400>  136

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1               5                   10


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1               5                   10


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<400>  138

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1               5                   10
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1   5   10

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1   5   10

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1   5   10

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1   5   10

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1   5   10

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1   5   10

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1               5                   10


<210>   147
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<400>   147

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1               5                   10


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<400>   148

Ile Arg Glu Asn Phe Lys Tyr Leu Gly Leu Trp
1               5                   10


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<400>   149

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1               5                   10


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1   5   10

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<400> 152

Phe Val Asn Cys
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Glu Val Arg Arg
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Arg Val Gln
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Arg Leu Thr Arg
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Ser Arg Lys
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<400> 157

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Met Leu Asn
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<400> 160

Leu Gln Arg Asn Lys Lys Gln Tyr Asn Leu Gly
1   5   10

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<400> 161

Leu Gln Leu Arg Tyr Lys Cys Tyr Asn Leu Val
1   5   10

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Val His Val Lys Leu Lys Glu Tyr Asn Leu Val
1   5   10

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<400> 163

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1   5   10

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1   5   10

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<400> 167

Gly Gly His
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<400> 168

Arg Ser Glu Val
1

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<400> 169

Cys Glu Pro Gly
1

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<400> 170

His Gln Met Ala
1

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Val Pro Ser Pro
1

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Gln Met Pro Asn
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Met Trp Pro Ser
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Cys Val Glu
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<400> 175

Leu Gln Leu Asn Leu Lys Val Tyr Asn Leu Val
1   5   10

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<400> 176

Leu Glu Leu Asn Leu Lys Val Tyr Asn Leu Phe
1   5   10

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<400> 177

Leu His Leu Asn Met Ala Glu Val Ser Leu Val
1   5   10

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<400>   178

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1                5                    10


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<400>   179

Pro Arg Gln Leu
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<400>   180

Phe Phe Trp Val
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Asn Phe Arg Asn
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<400> 183

Leu Gln Leu Lys Arg Gly Glu Tyr Ile Leu Val
1   5   10

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<400> 184

Cys Ser Leu Lys Leu Lys Ala Tyr Asn Leu Val
1   5   10

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<400> 185

Leu Gln Met Asn His Asn Glu Tyr Asn Leu Val
1   5   10

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Pro Gln Leu Asn Leu Asn Ala Tyr Asn Leu Val
1   5   10

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<400> 187

Gln Gly Met Gln Leu Arg Arg Phe Lys Leu Arg
1   5   10

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Arg Trp Leu His Trp Gln Tyr Arg Gly Arg Gly
1   5   10

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Pro Arg Pro Arg Leu Leu Arg Phe Lys Ile Pro
1   5   10

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Gln Gly Glu His Leu Arg Gln Leu Gln Leu Gln
1   5   10

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Gln Arg Leu Arg Leu Gly Pro Asp Glu Leu Phe
1   5   10

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<400> 192

Gln Arg Ile His Arg Arg Pro Phe Lys Phe Phe
1   5   10

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<400> 193

Gln Arg Met Pro Leu Arg Leu Phe Glu Phe Leu
1   5   10

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<400> 194

Gln Arg Val His Leu Arg Gln Asp Glu Leu Leu
1   5   10

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<400> 195

Asp Arg Met His Leu Trp Arg Phe Gly Leu Leu
1   5   10

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<400> 196

Gln Arg Met Pro Leu Arg Gln Tyr Glu Leu Leu
1   5   10

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<400> 197

Gln Trp Met Asp Leu Arg Gln His Glu Leu Leu
1   5   10

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<400> 198

Gln Arg Met Asn Leu Gly Pro Cys Gly Leu Leu
1   5   10

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<400> 199

Asn Cys Met Lys Phe Arg Ser Cys Gly Leu Phe
1   5   10

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<400> 200

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1   5   10

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<400> 201

His Arg Arg His Ile Gly Pro Phe Ala Leu Leu
1   5   10

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<400>  202

Glu Arg Leu His Arg Arg Leu Phe Gln Leu His
1               5               10


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<400>  203

Pro Cys Ile Gln Leu Gly Gln Tyr Glu Ser Phe
1               5               10


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<400>  204

Gln Arg Leu Arg Leu Arg Lys Tyr Arg Leu Phe
1               5               10


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<400>  205

Ile Val Glu Ile Leu Glu Asp Cys Gly Leu Phe
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<400> 206

Met Leu Asp Asn Leu Lys Ala Cys Gly Leu Phe
1   5   10

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Ile Leu Glu Asn Leu Lys Asp Cys Gly Leu Phe
1   5   10

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<400> 208

Leu Arg Glu Asn Leu Lys Asp Cys Gly Leu Leu
1   5   10

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Leu Leu Asp Ile Leu Lys Asp Cys Gly Leu Phe
1   5   10

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<400> 210

Val Arg Asp Ile Leu Lys Asp Cys Gly Leu Phe
1   5   10

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Ile Leu Glu Ser Leu Asn Glu Cys Gly Leu Phe
1   5   10

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Ile Leu Gln Asn Leu Lys Asp Cys Gly Leu Phe
1   5   10

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<400> 213

Met Leu Asp Asn Leu Lys Asp Cys Gly Leu Phe
1   5   10

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1   5   10

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1   5   10

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Ile Val Lys Asn Leu Glu Asp Cys Gly Leu Phe
1   5   10

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1   5   10

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Ile Arg Asp Asn Leu Lys Asp Cys Gly Leu Phe
1   5   10

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1   5   10

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<400> 220

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1   5   10

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1   5   10

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Pro Arg Asp Asn Thr Lys Val Arg Gly Leu Phe
1   5   10

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Phe Trp Gly Asn Leu Gln Asp Ser Gly Leu Phe
1   5   10

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<400> 224

Arg Arg Gly Asn Gly Lys Asp Cys Arg His Phe
1   5   10

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<400> 225

Leu Gln Glu Asn Leu Lys Glu Met Met Leu Gln
1   5   10

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<400> 226

Leu Glu Glu Asn Leu Lys Tyr Arg Met Leu Asp
1   5   10

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1   5   10

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<400> 228

Leu Gln Glu Thr Met Lys Asp Gln Ser Leu Gln
1   5   10

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<400> 230

Trp Gln His Lys Leu Ser Glu Val Met Leu Gln
1   5   10

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1   5   10

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Leu Leu Gly Met Leu Glu Pro Leu Met Glu Gln
1   5   10

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1   5   10

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1   5   10

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1   5   10

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Leu Gln Ala Asn Arg Lys Leu Gly Met Leu Gln
1   5   10

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Leu Ile Val Lys Val Lys Gln Leu Ile Trp Gln
1   5   10

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Met Arg Ala Lys Leu Asn Asn Leu Met Leu Glu
1   5   10

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Leu Gln Asp Asn Leu Arg His Leu Ile Gln
1   5   10

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<400>   240

Leu Arg Asp Asn Arg Asn Gln Leu Leu Phe
1   5   10

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1   5   10

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1   5   10

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Leu Gln Leu Asn Leu Lys Lys Tyr Asn Arg Val
1   5   10

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1   5   10

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1   5   10

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1   5   10

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Leu Gln Arg Asp His Val Glu Tyr Lys Leu Phe
1   5   10

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<400> 248

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1   5   10

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<400> 249

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1   5   10

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<400> 250

Met Gln Leu Asn Leu Lys Glu Tyr Asn Leu Val
1   5   10

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<400> 251

Val Gln Val Lys Leu Lys Glu Tyr Asn Leu Val
1   5   10

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<400> 252

Gln Leu Leu Asn Gln Tyr Val Tyr Asn Leu Val
1   5   10

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<400> 253

Trp Arg Leu Ser Leu Lys Val Tyr Asn Leu Val
1   5   10

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<400> 254

Leu Gln Arg Asn Lys Asn Gln Tyr Asn Leu Gly
1   5   10

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<400> 255

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1   5   10

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1   5   10

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<400> 257

Leu Phe Leu Asn Leu Lys Glu Tyr Ser Leu Val
1   5   10

<210> 258
<211> 11
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<213> Artificial Sequence

<220>
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<400> 258

Leu Glu Leu Asn Leu Lys Val Tyr Asn Leu Val
1   5   10

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Leu Pro Leu Asn Leu Ile Asp Phe Ser Leu Met
1   5   10

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Leu Pro Arg Asn Leu Lys Glu Tyr Asp Leu Gly
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Leu Arg Leu Asn Asp Ile Glu Ala Leu Leu Val
1   5   10

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Leu Val Leu Asn Arg Ile Glu Tyr Asn Leu Leu
1   5   10

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<400> 263

Leu Lys Arg Lys Leu Lys Glu Ser Asn Met Gly
1   5   10

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<212> PRT
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<400> 264

Leu Lys Arg Lys Val Lys Glu Tyr Asn Leu Gly
1   5   10

<210> 265
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<220>
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<222> (1)..(19)
<223> Reverse primer

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<210> 266
<211> 11
<212> PRT
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<220>
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<400> 266

Ile Leu Glu Asn Leu Lys Asp Cys Gly Leu Leu
1   5   10

<210> 267
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<212> DNA
<213> Homo sapiens

<400> 267
gccgccacc   9

<210> 268
<211> 57
<212> DNA
<213> Artificial Sequence

<220>
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<222> (1)..(57)
<223> Gi alpha 1/2 carboxy terminal sequence oligonucleotide

<400> 268
gatccgccgc caccatggga atcaagaaca acctgaagga ctgcggcctc ttctgaa   57

<210> 269
<211> 57
<212> DNA
<213> Artificial Sequence

<220>
<221> misc_feature
<222> (1)..(57)
<223> complementary strand to Gi alpha 1/2 oligonucleotide

<400> 269
agctttcaga agaggccgca gtccttcagg ttgttcttga ttcccatggt ggcggcg   57

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<211>  19
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<213>  Artificial Sequence

<220>
<221>  misc_feature
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<223>  forward primer for G alpha carboxyl terminal peptide insert


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atccgccgcc accatggga                                                    19


<210>  271
<211>  20
<212>  DNA
<213>  Artificial Sequence

<220>
<221>  misc_feature
<222>  (1)..(20)
<223>  reverse primer for G alpha carboxyl terminal peptide insert


<400>  271
gcgaaaggag cggggcgcta                                                   20
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